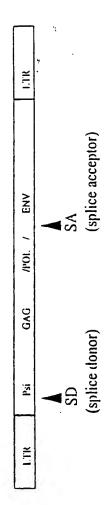
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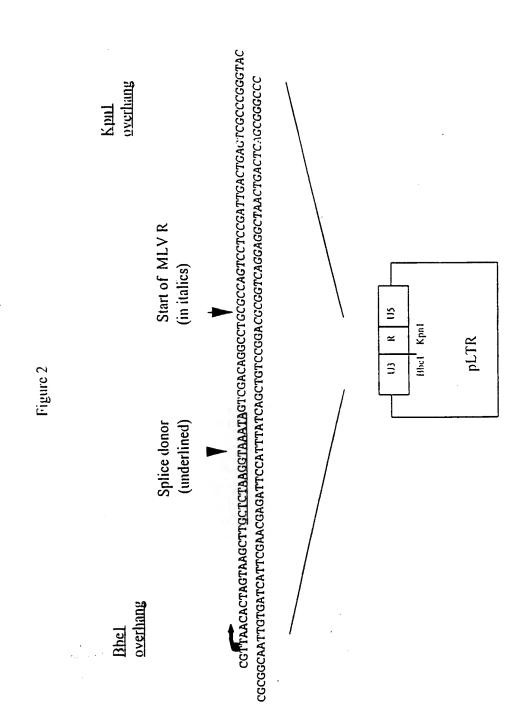
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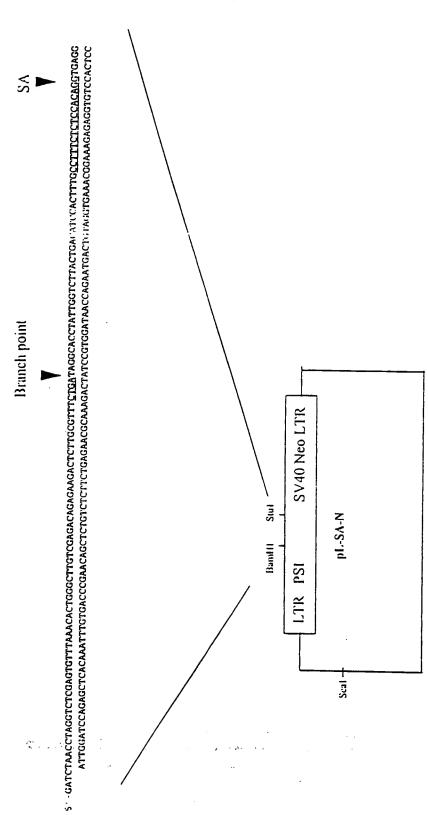
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Figure 3

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Figure 4

Spel U3 SD R U5 SA Neo Spel PSI GT TO GC CHANGE US R US Ascl 5 · · CCCTCACTCGGCGCGCCAGTCTTCCGA-3' Ascl

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GCTAKK TIAAGTAACGCCACTITIGCAAGGCATGGAAAAAATAKATAAAAATAGAAAAGTICAGATCAAGGTCAGGAACAAAGAAKAKKITGAATACCAAACAGGATATCTGTGGTAAAGG

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GACCCATGGCGATGCCTGCTTGCCGAATATCATGGTGGAAAATGGCCGCTTTTCTGGATTCATCGACTGTGGCCGGCTGTGGC;#GACCGCTATCAGGACATAGCGTTGGCTACCGTGA ataactgagaatagagaagttcagatcaaggtcaggaacagatggaacagctgaatatgggccaaacaggatatctgtggtaagcagificctggcccggctcagggccaagaacagatggaac IGCATCCGACTTGTGGTCTCGCTGTTCCTTGGGAGGGTCTCCTCTGAGTGATTGACTACCCGTCAGCGGGGTCTTTCA1111GGGG;CTCGGGGATCGGGAGACCCCTGCCCAGGGACCA ACAAGCCCGTCAGGGCGGTCAGCGGGTGTTGGCGGGTGTCGGGGCGCAGCCATGACCCAGTCACGTAGCGGAGTGTATAC IGGCTTAACTATGCGGCATCAGAGCAGATTGTACTG AGAGTGCACCATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCCGCTTCCTCGCTCACTGTGCGCTCGGTCGTTCGGCTGCGGCGGCGAGCG GINTCAGCTCACTCAAAGGCGGTAATACGGTTATCCACAGAGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTTGCTGGCG A THARCAGE CONGESTATION OF CHOLONG THE TITLE THE ANGLES OF TACES OF TACABAGGACAGTATING THE CHOCGEST OF TACTED AND THE CONTRACT OF TACES AND THE CONT GIST.TICTIGACGCTCAGTGCGAACGAAAACTCACGTTAAGGGTTTTTGGTCATGAGATTTTTCAAAAAAGGATCTTCACCTAGATCCTTTTAAAATTAAAATGAAATTTTAAATTTAAATTTAAATTTAAATTTAAATTTAAATT GAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTCCACGGCTCCAGATTTATCAGCAATAAACCAGCCAACCAGCCGAGCGCAGGGCGCAGAAGTGGTCCTGCAACT TTATCCGCCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAGTAGTTCGCCAGTTAATAGTTTGCGCAACGTTGTTGCT ATTGCTGCTGCAGGCATCGTGGTGTCACGCTCGTCGTTGTTT GGTATGGCTTCATTCAGCTCCGGTTCCCAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTTV GGTCGTTGTCAATGGTAGTAAGTTGGCCGCA GIGITPATCACTCATGGTTATGGCAGCACTGCATAAITICTIACTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGATACTGAGTACAAGTCATTCTGAGAATAGTGTATGCGGCGA CCGAGTTGCTCTTGCCCCGGCGTCAACACGGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGCTCATTGGAAAACGTTCTTC\\GGGCGAAAACTCTCAAGGATCTTACCGCTGTTG acacggaaatgttgaatactcatactcttcctttttcaatattaatgaaggattattgagggttattgtcicatgagcggatacatgtaatagtatttagaaaaattaagaaaatt CCGCGCACATTTCCCCGAAAAGTGCCACCTGACGTCTAAGAAACCATTATCATGACATTAACCTATAAAAATAGGCGTATCAACACATTCGTCTTCAAGAATTCATACCAGATCAC ITKKGGTTCKGGAGUTCAATAAAAGAGCCCACAACUKCTCAGTKGGCKJCGCCAGTCTTCCGATAGACTGCGTCGCCCGGGTACCCGTATTKAAAGCCTCTTGCTGTTTGCATCCGAAAT CGNGTCTCGCTGTTCCTTGGGAGGGTCTCCTCTGAGTGATTGACTACCCACGACGGGGGTCTTTCATTTGGGGGCTCGTCCGGGATTTGGAGACCCCTGCCCAGGGACCACGACCACCACCAC AGTYITTAAACACTGGGCTTGTCGAGACAGAGAGACTYYTTGCGTTTTYGTAGGCACCTATTGGTCTTACTGACATCCACTTTGCCTTTCCACAGGTGAGGCCTAGGCTTTTGCAAAA GUCA GUG C G G C TATO GT G G C C C C G G G G G G C T T G C G G G C T T G T C C T G A A G C G G G A A G G G C T TATIGGIGAAGAGCTIGGGGGGGAATGGGGCIGACCGCITCGTGGTTTACGGTATCGCCGCTCCCGATICGCAGCGCATCGCCTTV:TATCGCCTTCTTGACGAGTTCTTGAGGGGGACT agctigaatatigggccaaacaggatatutgttggttaagcagtttgcuccggctcagggccaagaacagatggtccccagatguugtuugtctagcagtttctagagaaccatcagattgtt TTTFICCATAGGCTCCGCCCCTGACGAGCATCACAAAATCGACGCCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGC GITICETOCCCGGCTCAGGGCCAAGAACAGATGAGACAGCTGAGGGTGATGAGGGATATCTGTGGTAAGCAGTTCCTGCCCCGGGCCCAAGAACAGATGGTCCCCAAGATG ICCEAC CCTCAGCAGITTCHAGTGAATCATCAGATGITITCCAAGGACCTGAAAGAACCTGAAATGACCCITGITACTITGAACHAATCAGITCGCITCTCGCTTCTGTTCGCGCGC GGAACTGACGAGTTCTGAACACCCGGCCGCAACCCTGGGAGGTCCCCAGGGACTTTGGGGGGCCGTTTTTGTGGCCCGACCTGAGGAAGGGAGTCGATGTGGAATCCGACCCCGTCAGGATAT GTKKTTKTRGTAGGAGAAGGAAGAAGTTCCCGGCTCCGTTTGAATTTTTGCGTTTGGAACCGGAAGCCGCGTCTTGTTGGAAGCCGCGTCTTGTGTGGAAGCGCGCTGCAGCATCGTTCTGTGTTGT HGACHGGGCACAACAGACAATCGGCTGCTCTGATGCCGGCGGCGCTGTTCCKHGCTGTCAGCGCGCGCGCGGGTTCTTTTGTCAAGACHGACHGTGTCCGGTGCCCTGAATGAACHGCAGGACGA TACTCGGATGGAAGCCGGTCTTGTCGATCAGGATGATCTGGACGAAGAGCATCAGGGGCTCGCGCCAGCCGAACTGTTCGCCAGGCTCAAGGCGCGCATGCCCGACGGCGAGGATCTCGTCGT TECAGGGTGCCCCAAGGACCTGAAATGACCCTGTGCCTTATTTGAACTAACCAATCAGTTCGCTTCTGGCTTCGCGCGCTT(`TGCTCCCGAGGTCAAAAAGAGCCCACAAACAACCCCT CACTCGGGGCCCGTTAACACTAGTAAGCTTGCTCTAAAGTAATATGTCGACAGGCCTGCGCCAGTCCTCCGATTGACTGAGTCGCCGGGTACCCGTGTATCCAATAAACCCTCTTGCAGT CTGGGCTGTGTGCGCCGCGCTTCAGCCCGAGCGCTGATCGCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGCCGCTTATCGCCACTGGCAGCAGCAGCAGTGGTAACAG SCITTIGAAAGACCCCACCCGTAGGTGGCAA

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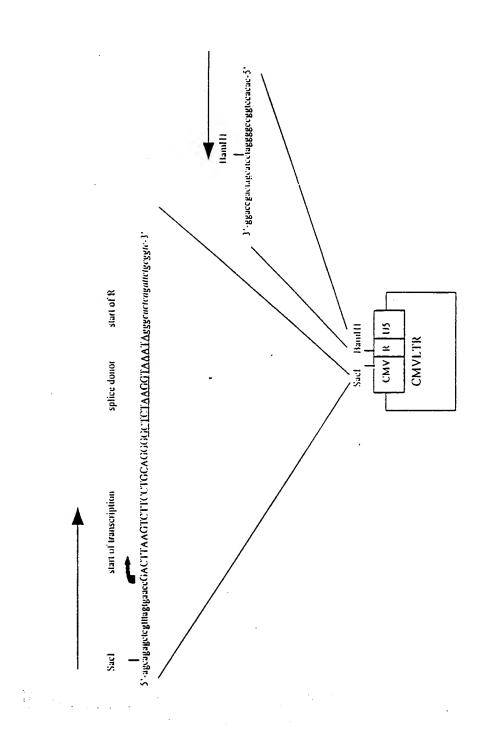


Figure 6

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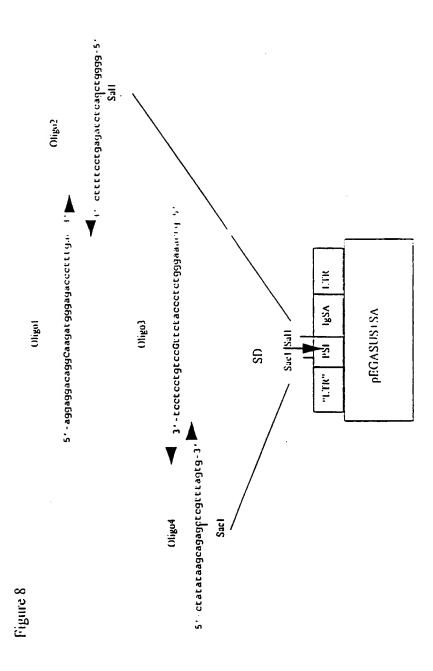
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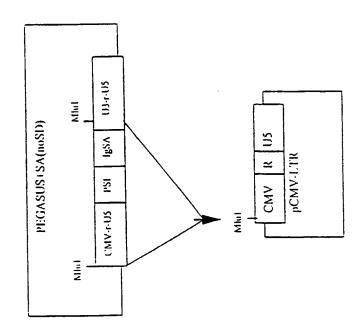
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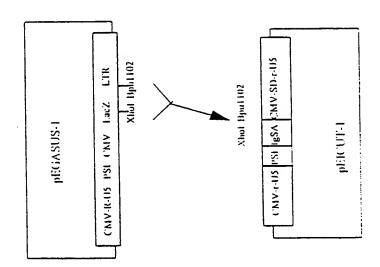
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Figure 11

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(A) pICUT vector in transfected cells

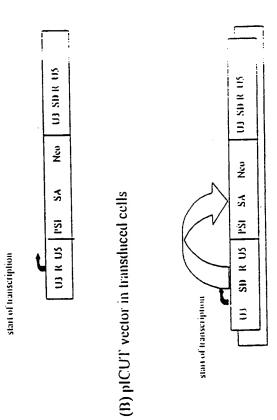
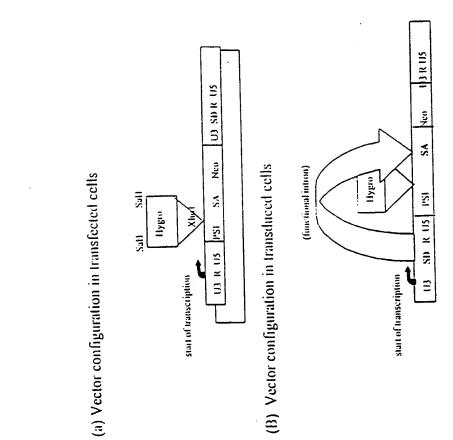


Figure 13



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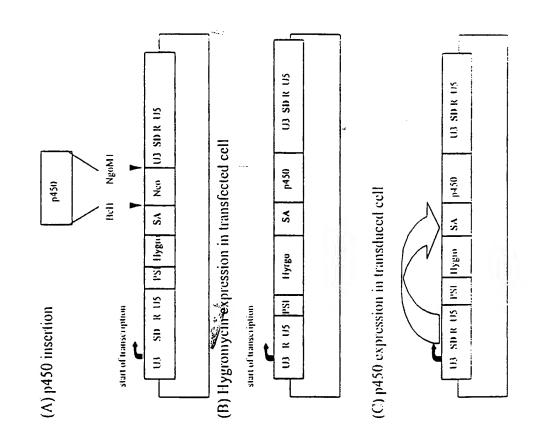


Figure 14

5' -ATG CGT TCA ACG CTC TCA AAA CCC CTT AAA AAT AAG 5' -ATG GCC AGA AGC ACC CTG AGC AAG CTA CCC CAG GAC 5'altered 4070A 3'end of pol

Figure 15

GTT AAC CCG CGA GGC CCC CTA ATC CCC-3'

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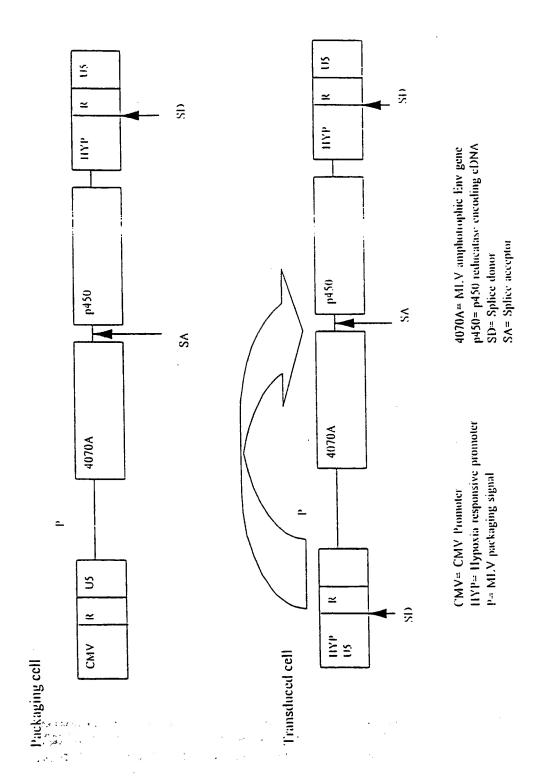
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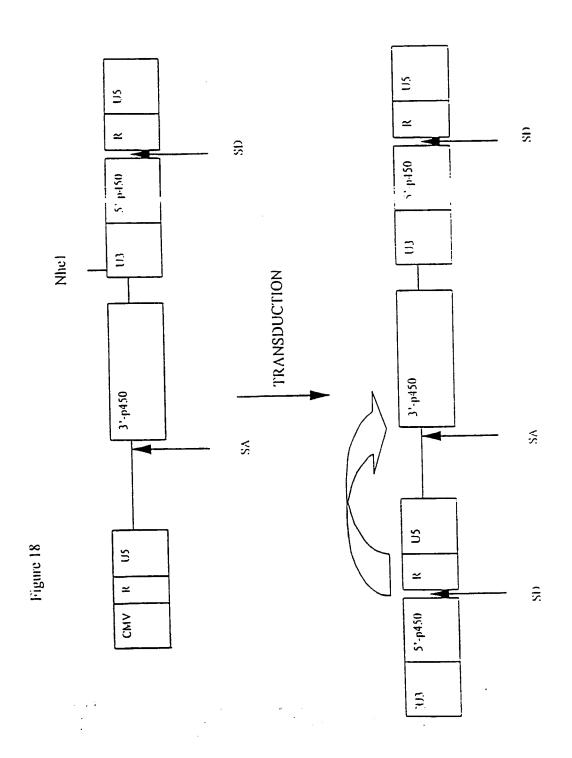
TIGGCACCTCC CTCCTGGGAA CTGTACAAGA TGUCTTUUTA IGTGGCCCCT GCTACGACCT CICCAAAGTA TCCAATTCCT TCCAAGGGGC TACTCGAAAA GGAGAGGAGT GGGACCCTTC AGACCAGGAA CCGTATGTCG GGTATGGCTG CAAGTACCCC GCAGGGAGAC AGCGGACCCG GACTTTTGAC TCGGGGTGTG GGGGACCAGG AGAGGGCTAC GGAAGCCCAC ATCATCGTHG TAAAGITIGE CCATAGGGCC CAACCCAGTA CTCCCCTA:JA TCTAGTCAAA ACAGGACAGG GCCTATGCAT GGGGCAGTA CCTAAAACTC ACCAGGCCTT ATGTAACAAG TAACTUGGAC TACCATGITTC ACAGCCACT Trecenerati TGGGCACTTA TACCAATCAT ATCTGAAGTG AATGTGGGGT CATTGACCCT GGCCCTTCTA CACAGATTAT CCCCCGA1TA TATGTATGTT CACTGCCTTA CAACGAAGTEC AGACCACACG GGGCTAGTGA GAGACAGCAT GGCCAAATTA ACAAAAACTA TTTGAGACAG GCCAAGGATG GTTCGAAAAA GITTACCACC ITAAICICCA CCATCAIGGG ACCICIAAIA PUTCACIOGT CCAGGUICIG GTITTGACTC AGGAATATG CCAGUTAAAA GACCTAATCT CCCTTAAGCG CGGTAACACC CCCTGGGACA CGGGATGCTC FIRECCUACC ANAGACTECE TITECTERECA ATAGAGATTG TACEGGETEC CACCCTCAAC CCCAAGAATG FCCACCGCTC GGCCAACTG TACGGCCACT TCCCAACATA AGCTTACCCT ACTIGATGCAG GAAAAAGGC CAGATCCTAT GACTACTAGC CCGCCGGAAC TCAATCTAAC GGACGGGGAC AGACAGACCT AGAAAAGTCA CTGACCTCGT TGTCTGAAGT IGTGGTAAAT GGGGGTGTGA AACCACCGGA CAGGCTTACT PCCCTGACCC GGCAGGTCCT TAATGTGGGA CCCCGAGTCC AGCCCCTCA ATACCAGTTA CCCCCTTCC ACTACCAGTA CCCGACAAGA TTAGTGTCGG GACCTCCTTA TTACGAAGGA GTAGCGGTCG AAGGAGGGAG CGGACAGGAA AGTICCAAGTG TICCCACAGCC ACCCCCAGGA ACTGGAGATA ACCCAAAGCG CCGGCTCAGG ATCCTACTAC CTTGCAGCAC ACCACGGTGC GCCCAGAGTA ATTIACCACT CAGCTIGAAC AGCGIACCAA ATATAAAAGA GAGCCAGTAT GCTGGAATAG GCCGCTATCC TIGATCTATG TGATCTGGTC I'TT'TACGTGT GCCCTGGGCA TACCGTAAAG ACCCTCTAGT CCTAGAATTC CGTGGGGACT GAGACTGTAC CCTCACCAAT CTGCTTGTCC CTAGGAGGAT TAACCATGGG AGGGATTGCA GCAGCTTCAT GCTATTCCTA GTACCGCCAA CCTGGAGAGT GGAGCCTATC AGGCGCTTAA TGCAGCACTG GATTGACTCC TGTGTATTAG TEGAACTCTG GAAAAGTCAA TTACCAACCT SAAGAATGTT GTTTTTATGC NGAGAAAGGC TTAATCAGAG CTGTTTAATA GATCCCCCTG CCATAGAGT ACGAGCCATG ATTAMAACCC AGCAGITTGA AACCGCAGAG GCCTAGATTT ATGACTGGGC AAATTATAT"F GGCAGATGCA GGCCCCAAAT ATCGTCATGG I.T.FAATGTAA

Figure 17

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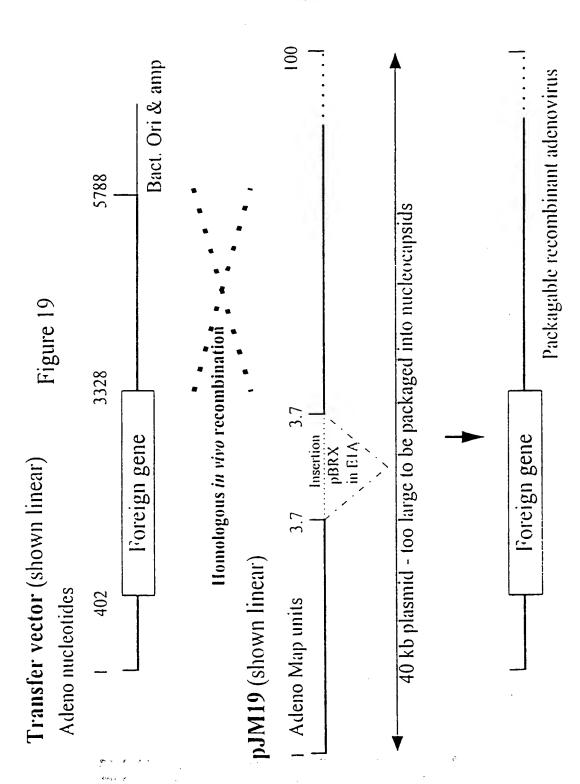


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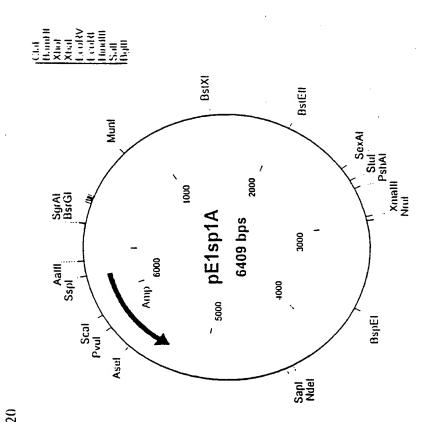
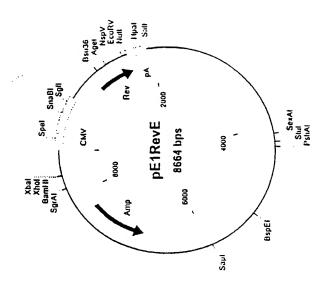


Figure 20



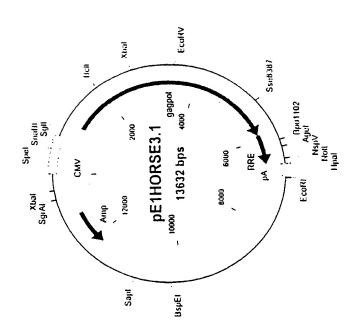
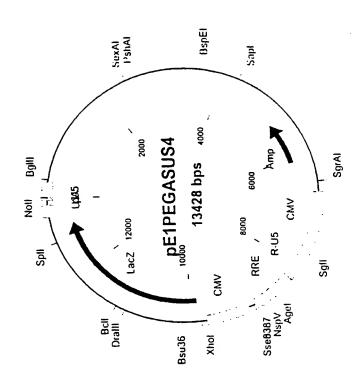
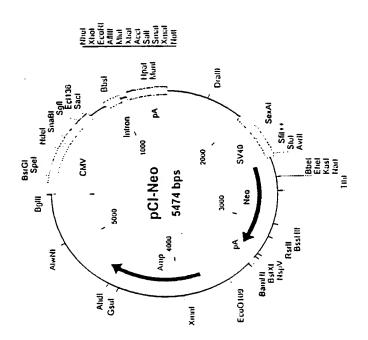


Figure 22



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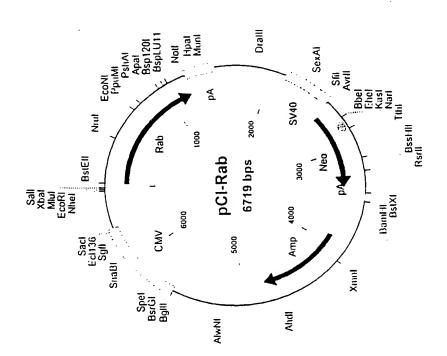
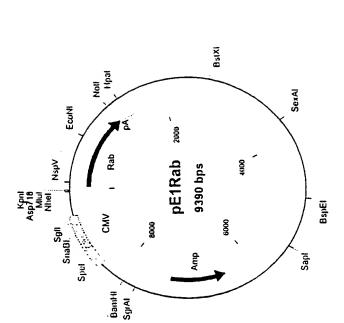
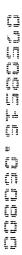
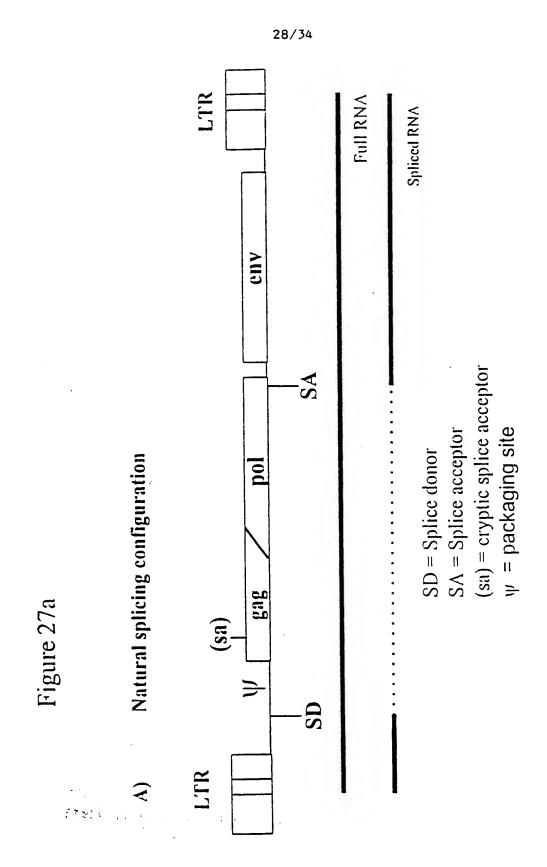


Figure 2



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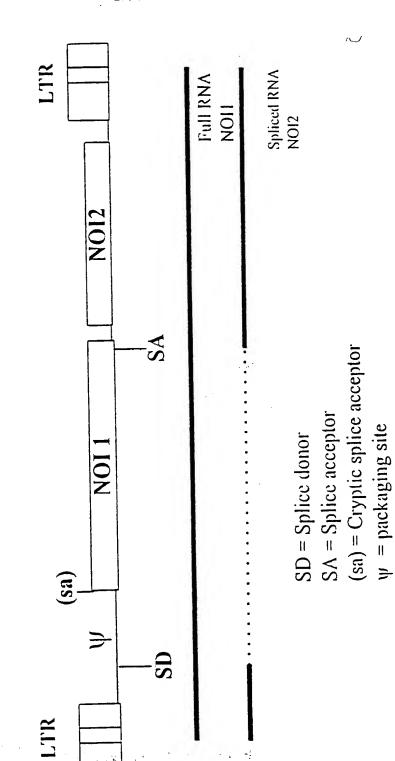
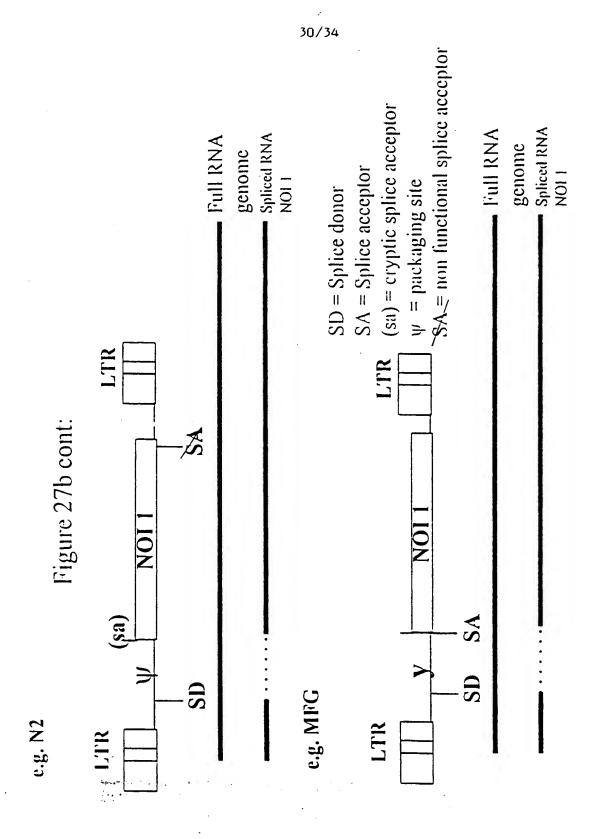


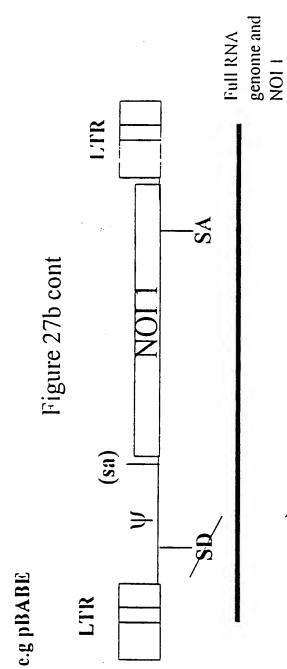
Figure 27b

Splicing configurations in known vectors

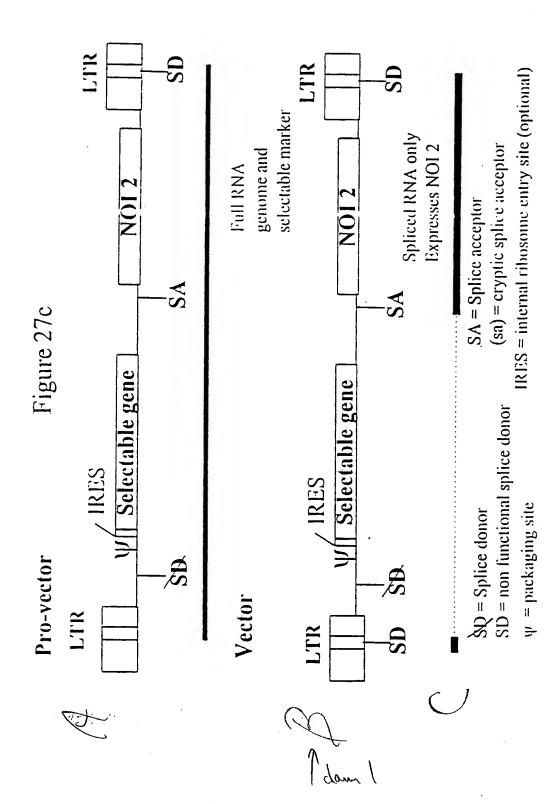
e.g. LTRSVX

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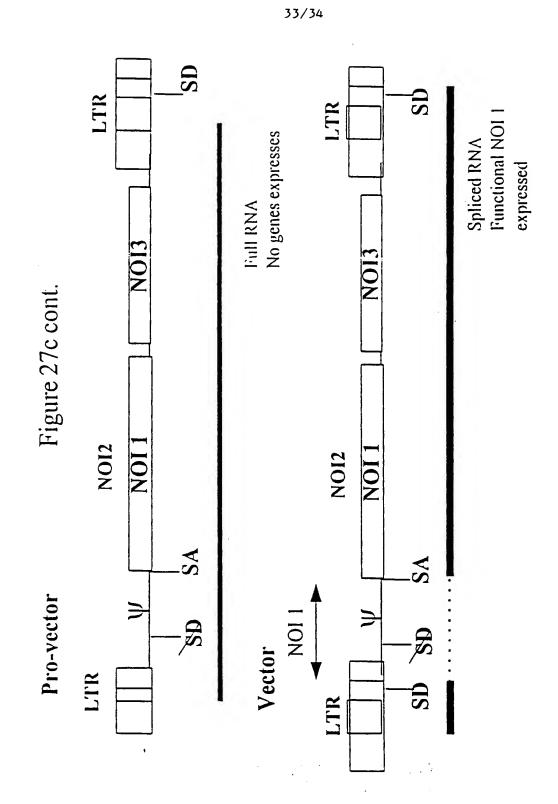




SR = Non functional splice donor SA = Splice acceptor (sa) = cryptic splice acceptor ψ = packaging site



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